

INFLUENCE OF GEOMETRICAL NUCLEAR PROPERTIES ON CHROMOSOME ABERRATION

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Background

- **Space radiation.** Among main hazards for space flight, primarily due to galactic cosmic ray high charge and energy (HZE) ions.
- **Relative biological effectiveness (RBE).** Important factor used to estimate space-radiation quality factor in cancer risk models [1].
- **Chromosome aberrations (CA).** Possible biomarker to assess HZE ion RBE and related cancer risks [2].
- Role of nuclear 3-dimensional (3D) genomic organization on the formation of CAs remains to be understood [3].

Objectives

- To assess how **change in nuclear properties and chromosome organization impacts CA formation in terms of RBE values.**
- To **compare to experimental results** for heavy ions of linear energy transfer (LET) relevant to space radiation [4-7].

Material & Methods

- 46 interphase chromosomes modelled as monomer sequences distributed in the nucleus with a random walk. Chromosomes associated with non-overlapping domains.
- Two chromosome intermingling configurations:
 - **Min.** Monomer sequence constrained in domains.
 - **Max.** Only first monomer of sequence constrained in domains.
- Cell nuclei geometries
 - Spherical with radii R varying from 2 μm to 8 μm .
 - Ellipsoidal with volume $V = 162 \mu\text{m}^3$ (fibroblast AG1522) or $V = 656 \mu\text{m}^3$ (fibroblast 82-6) and thicknesses z varying from 0.5 μm to 5.38 μm .
- Monte-Carlo track-structure tool **RITRACKS** [8] used for **transport of HZE ions and γ -rays** (0.6617 MeV) in cell nuclei, for doses D in the range 0 to 1 Gy.
- Monte-Carlo tool **RITCARD** [9,10] used to model DNA double-strand breaks (DSB) (\sim 35 DSB/Gy/cell), DSB (mis)-repair for a 24h time-period and **chromosome aberration classification**.
- Linear quadratic dose response for **total exchange**
$$y(D) = \alpha D + \beta D^2 \text{ and } RBE = \alpha_{\text{HZE}}/\alpha_{\gamma}$$

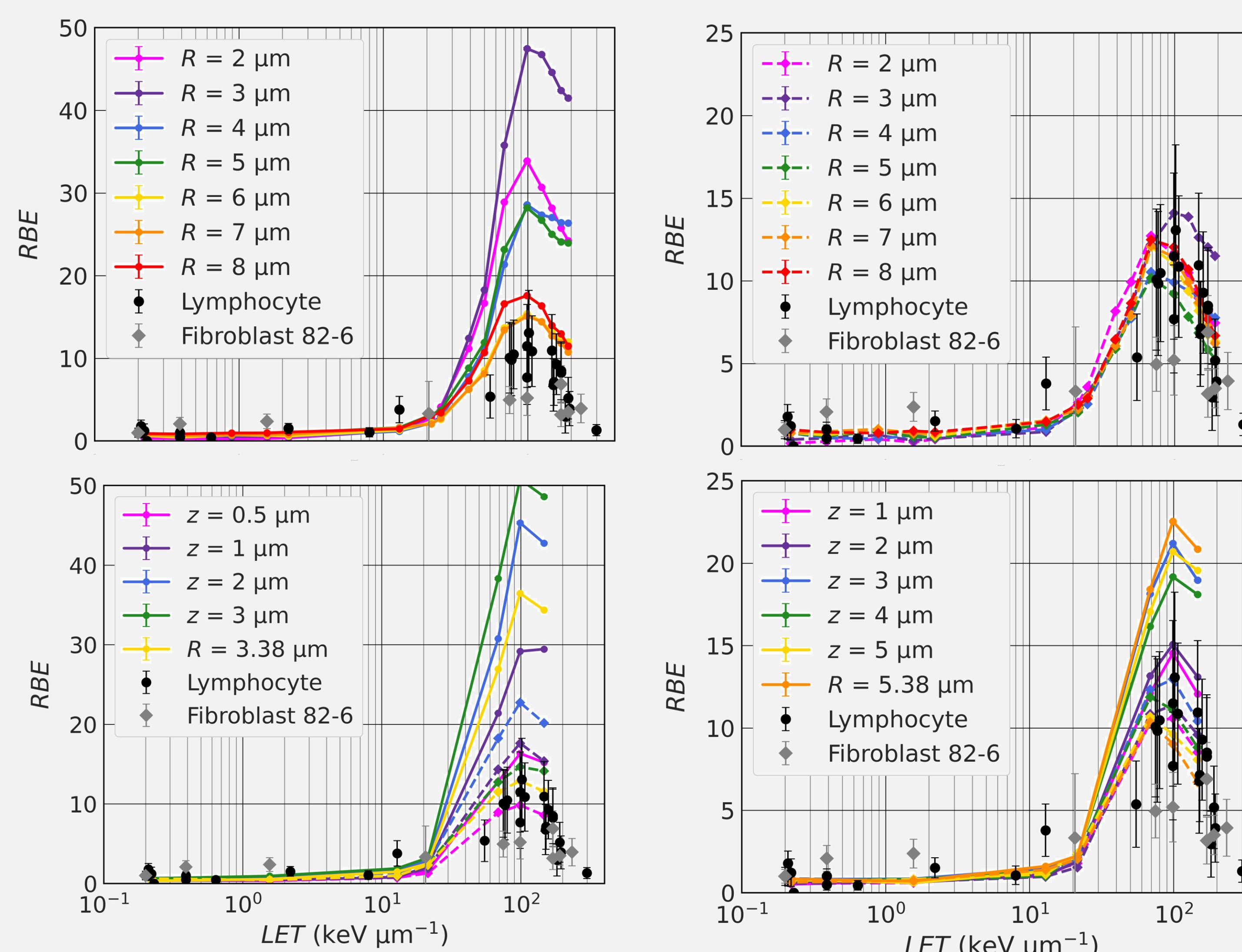
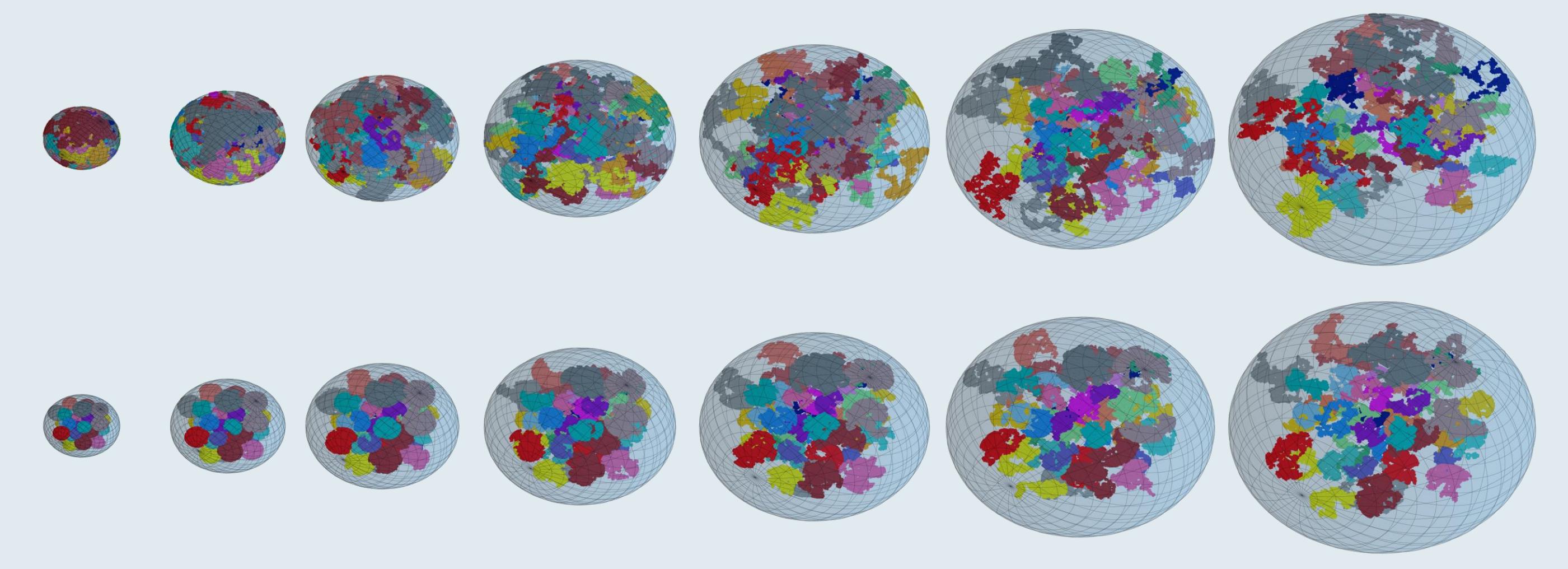
References

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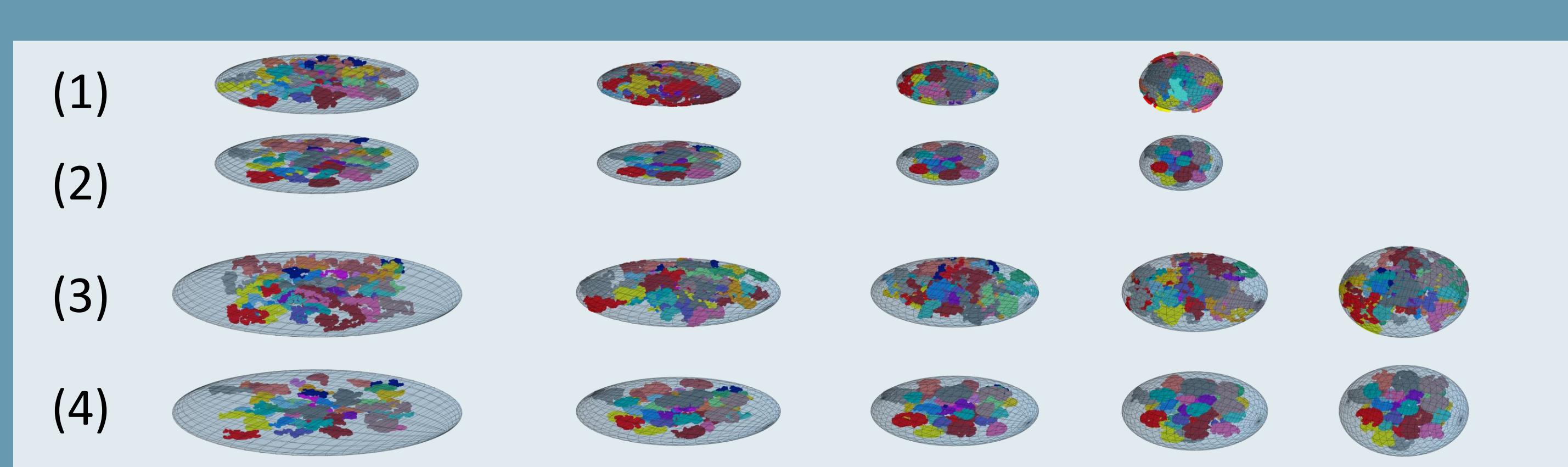


Results & Discussion

Chromosome distribution in spherical nuclei with radii increasing from 2 μm to 8 μm . Top: maximum intermingling. Bottom: minimum intermingling.



RBE as a function of LET for total exchanges. Exp. data from [4-7]. Top: Spherical. Bottom: Ellipsoidal; left: AG1522 ($V = 162 \mu\text{m}^3$), right: 82-6 ($V = 656 \mu\text{m}^3$). Solid: max. intermingling. Dashed: min. intermingling.



Chromosome distribution in ellipsoidal nuclei. (1) & (2) $V = 162 \mu\text{m}^3$, $z \in 0.5 \mu\text{m}$ to 3 μm . (3) & (4) $V = 656 \mu\text{m}^3$, $z \in 1 \mu\text{m}$ to 5.38 μm . (1) & (3) Maximum intermingling. (2) & (4): minimum intermingling.

➤ LET effect

- RBE peaks at $\sim 100 \text{ keV}/\mu\text{m}$ regardless of size, shape or chromosome intermingling, consistent with experimental data.
- Little dependence on 3D genomic organization for low-LET ions.

➤ Size effect

- RBE dependence on R for high-LET ions and max. intermingling, with increase of RBE from 2 μm to 3 μm , followed by a decrease of RBE from 3 μm to 6 μm and saturation beyond.
- Little RBE dependence on R for min. intermingling.

➤ Shape effect

- AG1522 (small nucleus). RBE dependence on z , with increase of RBE from $z = 0.5 \mu\text{m}$ to 3 μm and decrease for spherical nucleus for max. intermingling. Weaker dependence on z for min. intermingling, with RBE peaking for $z = 2 \mu\text{m}$.
- 82-6 (large nucleus). RBE dependence on z to a lesser extent compared to AG1522. Increase and saturation of RBE with increasing z for max. intermingling. Little dependence on z for min. intermingling.

➤ Min. vs. Max. Intermingling

- Decrease of RBE values when intermingling reduced.
- Decrease of RBE dependence on nuclear geometry when intermingling reduced.

➤ Lymphocyte vs. Fibroblast

- Lymphocyte. Spherical, $R \sim 3 \mu\text{m}$, $V \sim 113 \mu\text{m}^3$.
- 82-6. Ellipsoidal, $x \sim 7.22 \mu\text{m}$, $z \sim 3 \mu\text{m}$, $V = 656 \mu\text{m}^3$.
- Higher RBE values for lymphocyte vs. fibroblast, consistent with geometrical effect.
- Comparable RBE values between experimental data and simulated results for min. intermingling.

Conclusions

- RBE dependence on size and shape of the nucleus, with effects more pronounced with max. intermingling.
- Trends due to dependence of α_{HZE} and α_{γ} on 3D genomic organization for both simple and complex exchanges [11].
- Nuclear geometrical properties to be assessed when comparing CA between cell lines of different size and shape, or when translating *in vitro* results to *in vivo* risks.

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